

# Site – Specific Monitoring For Diseases Forecasting in Winter Wheat

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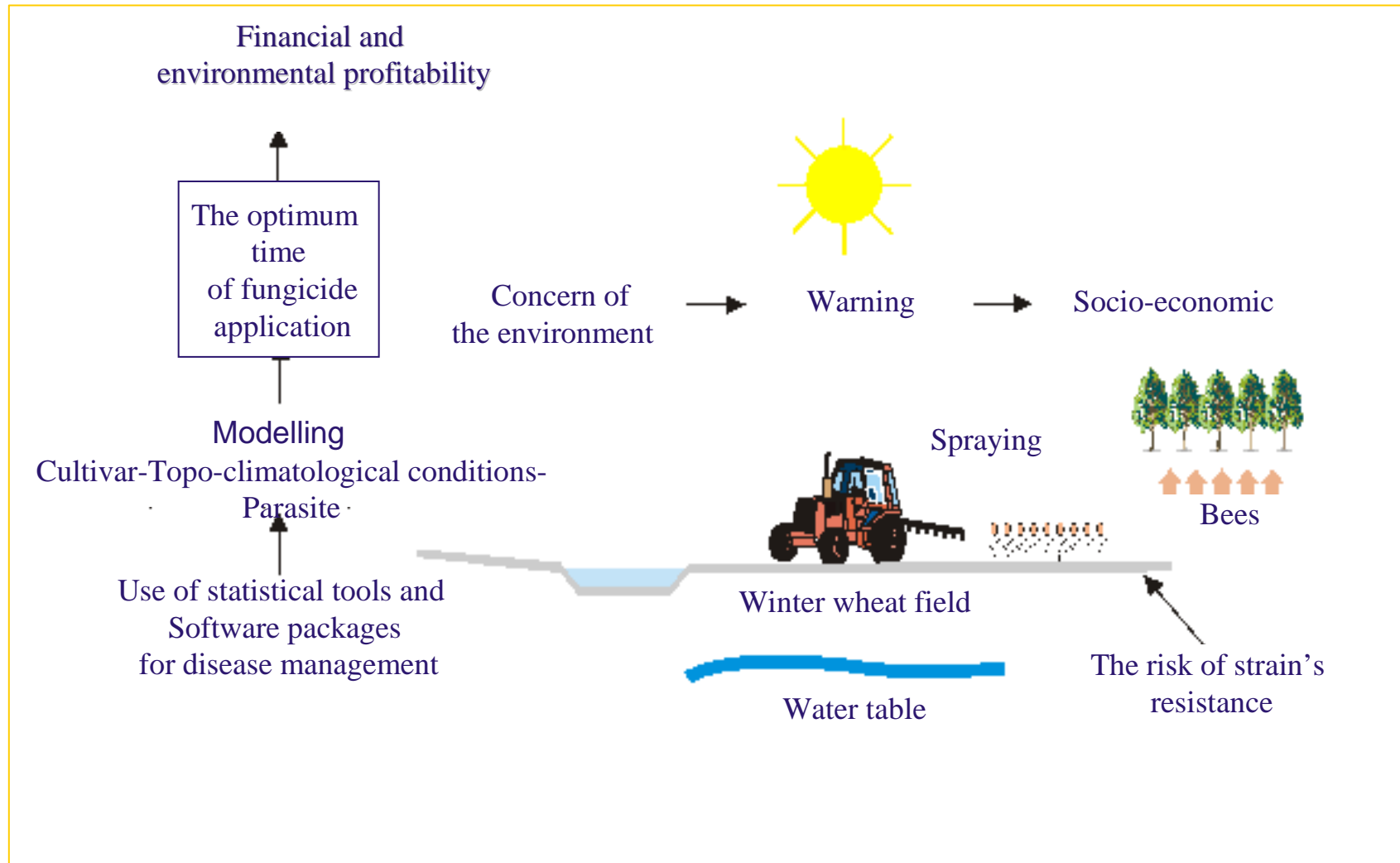
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# Introduction

## Why to set up a decision support system for disease control in wheat?



# Introduction



Powdery mildew



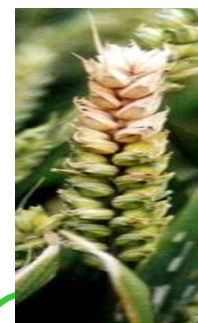
Septoria



Yellow rust

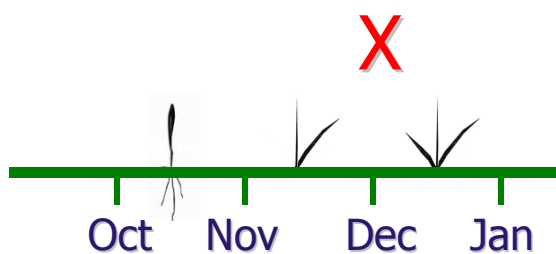


Leaf rust

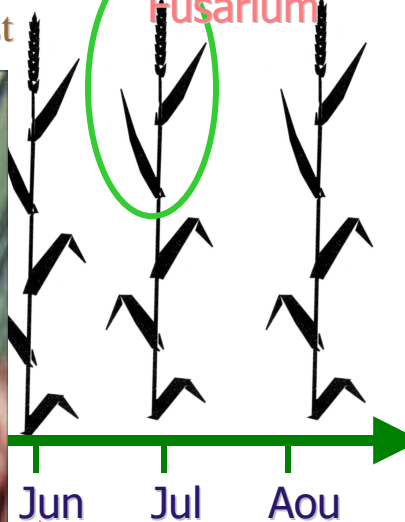


Fusarium

The goal is therefore to provide  
frame for site-specific spraying  
setup for sites throughout Luxem



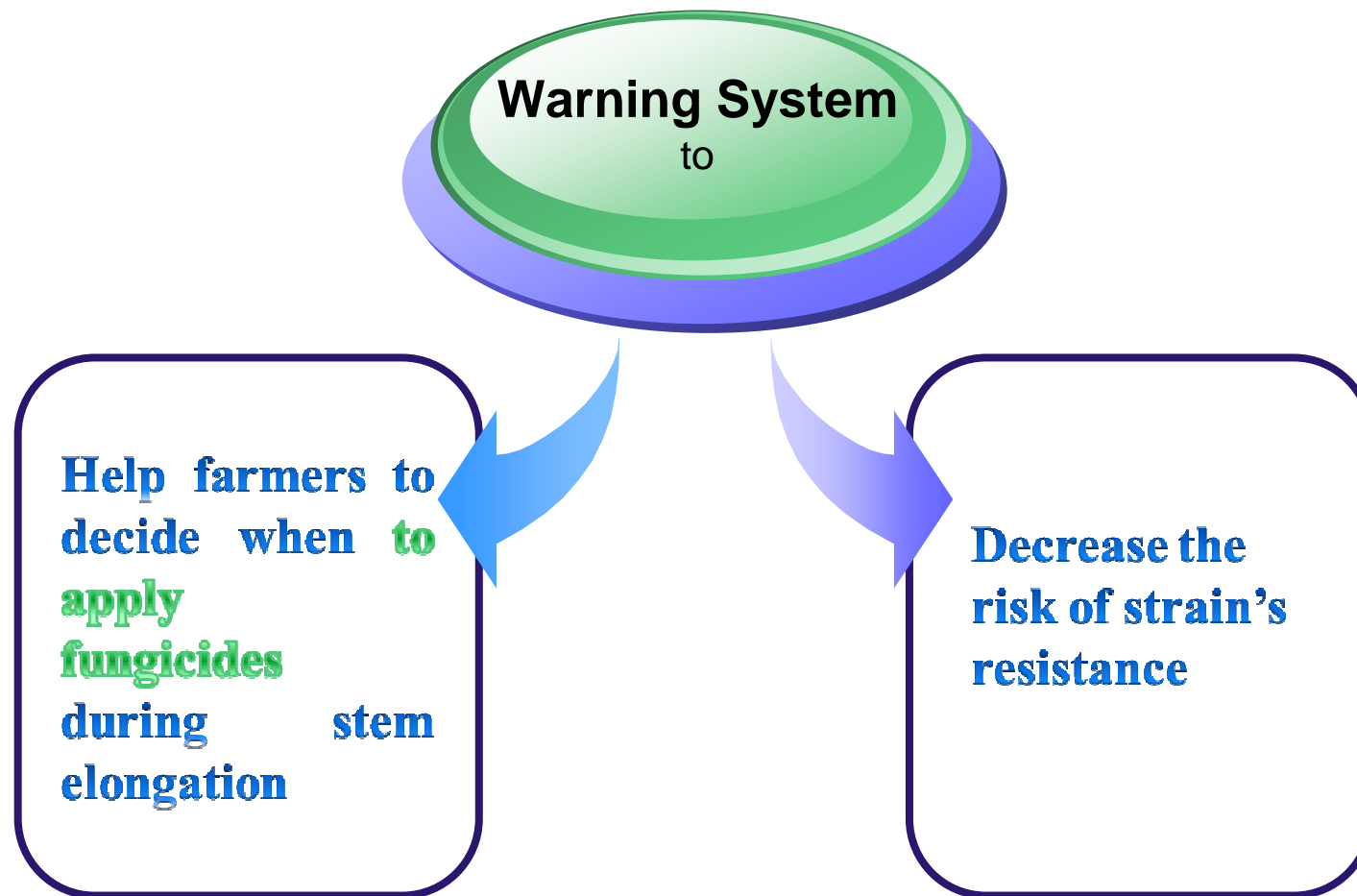
Objectif



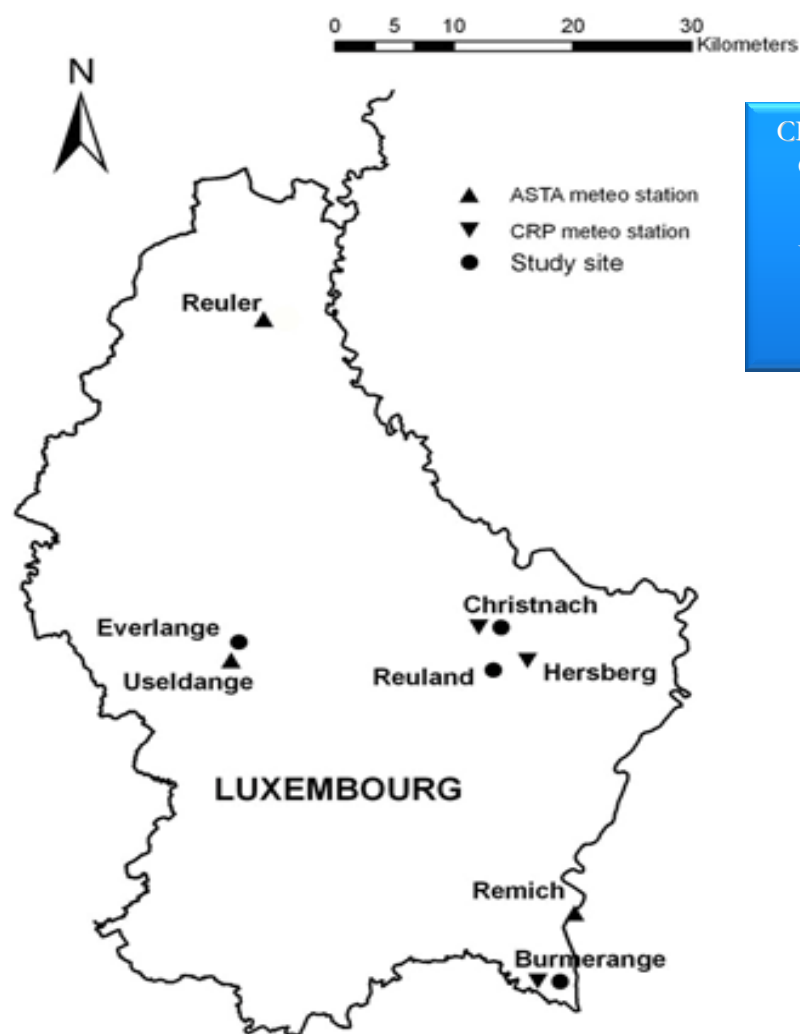
Average: 1,6

Generally  $\leq 1$

# Objectives of the work



# Methodology



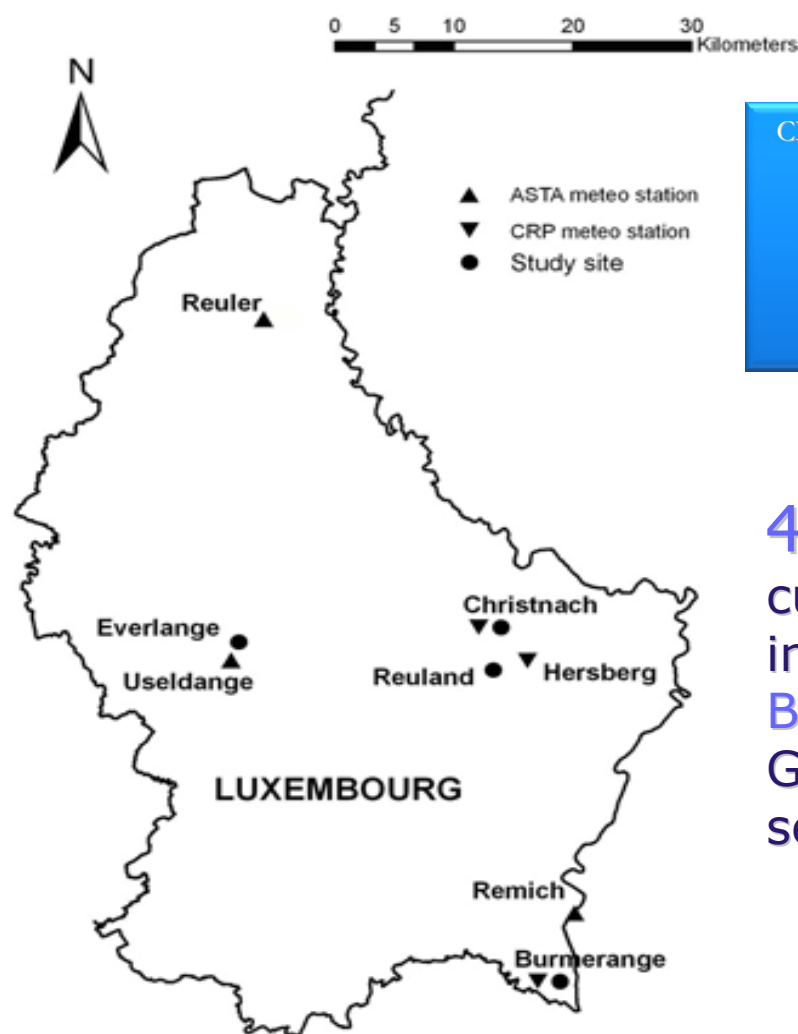
CRP: Public Research Center  
 Gabriel Lippmann, for its  
 initials in French  
 ASTA: Administration of  
 Agricultural Technical  
 Services, for its initial in  
 French.



# Methodology



# Methodology

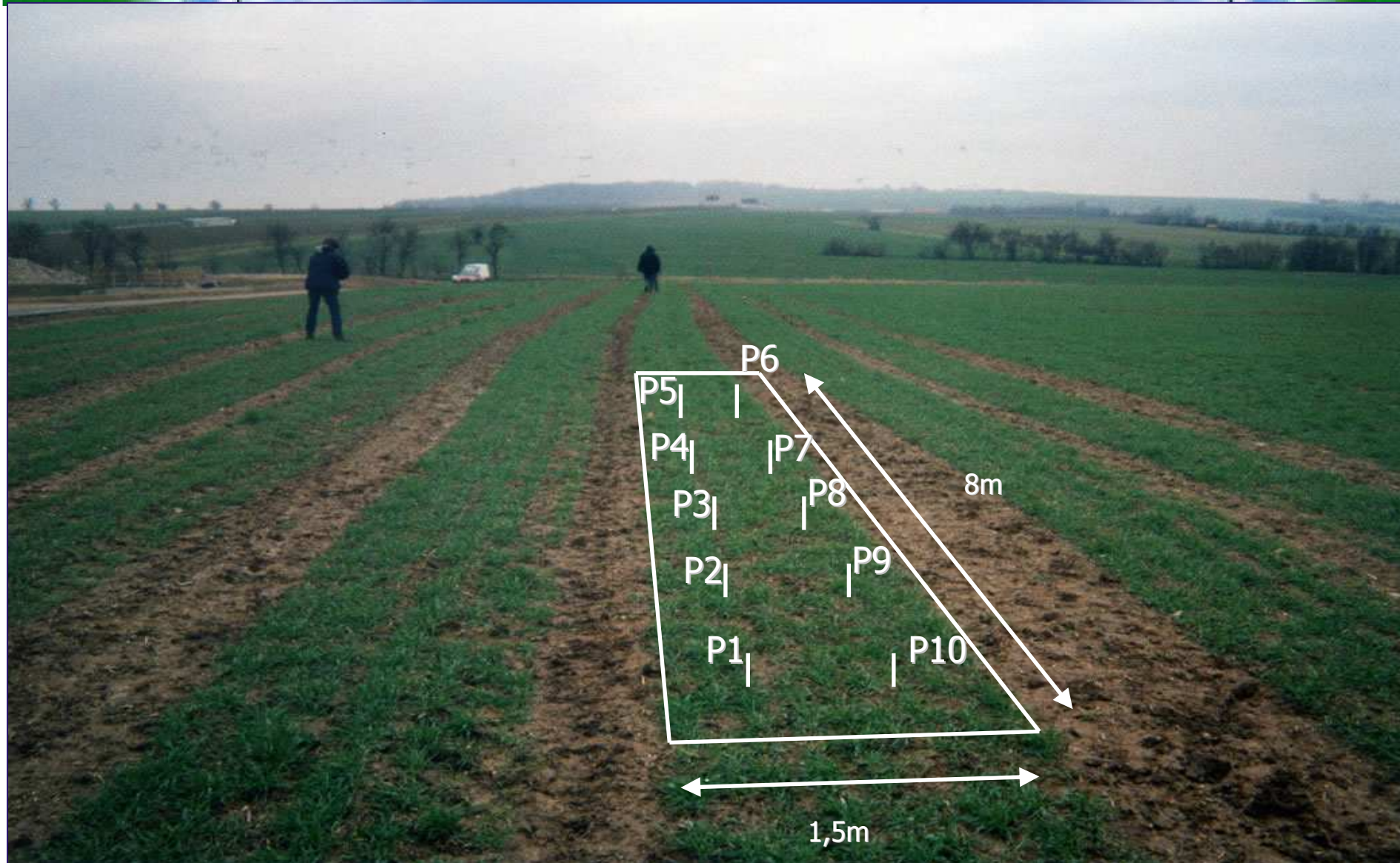


CRP: Public Research Center  
Gabriel Lippmann, for its  
initials in French  
ASTA: Administration of  
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Services, for its initial in  
French.

4 replicated field experiments on 2 cultivars (minimum) were established in the sites of Everlange, Burmerange, Christnatch and Reuler G.D. of Luxembourg, for the growing seasons for 11 years.



# Methodology



# Methodology



Standard area diagrams of James (1976) and  
software programs DISTRAIN



# Septoria Leaf Blotch <sup>1/4</sup>

Logarithmic scale

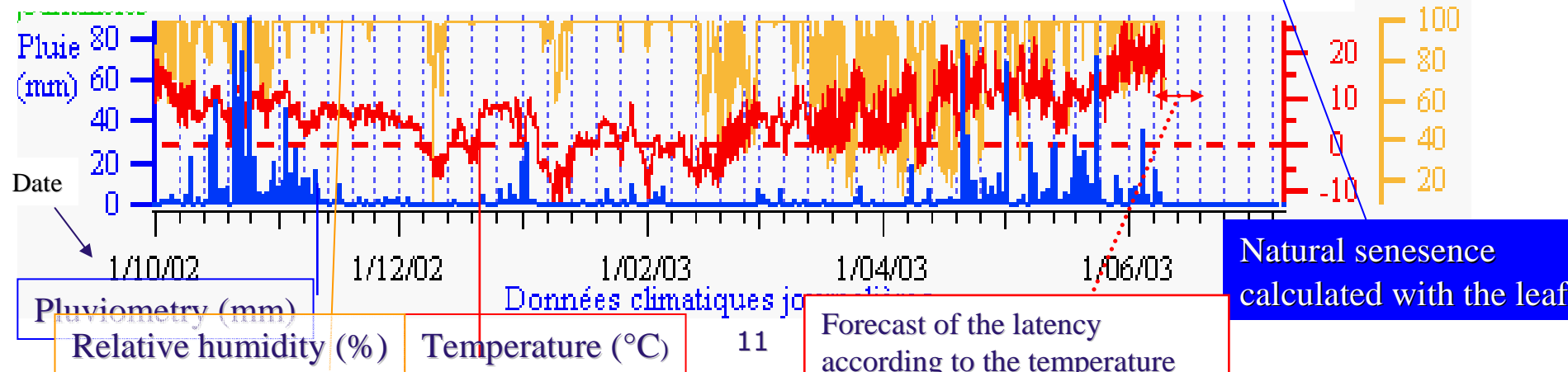
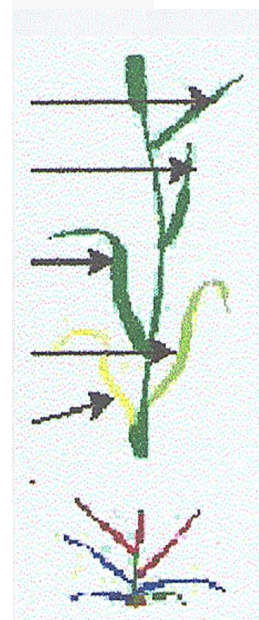
Percentage of the leaf area covered by sporulating *S. tritici*  
(Pink: Primary infection, red: secondary infection)

Forecast of the percentage of the area covered by sporulating *S. tritici*  
(beige: Primary infection, brown: Secondary infection)

Leaf area development

Example of latent period

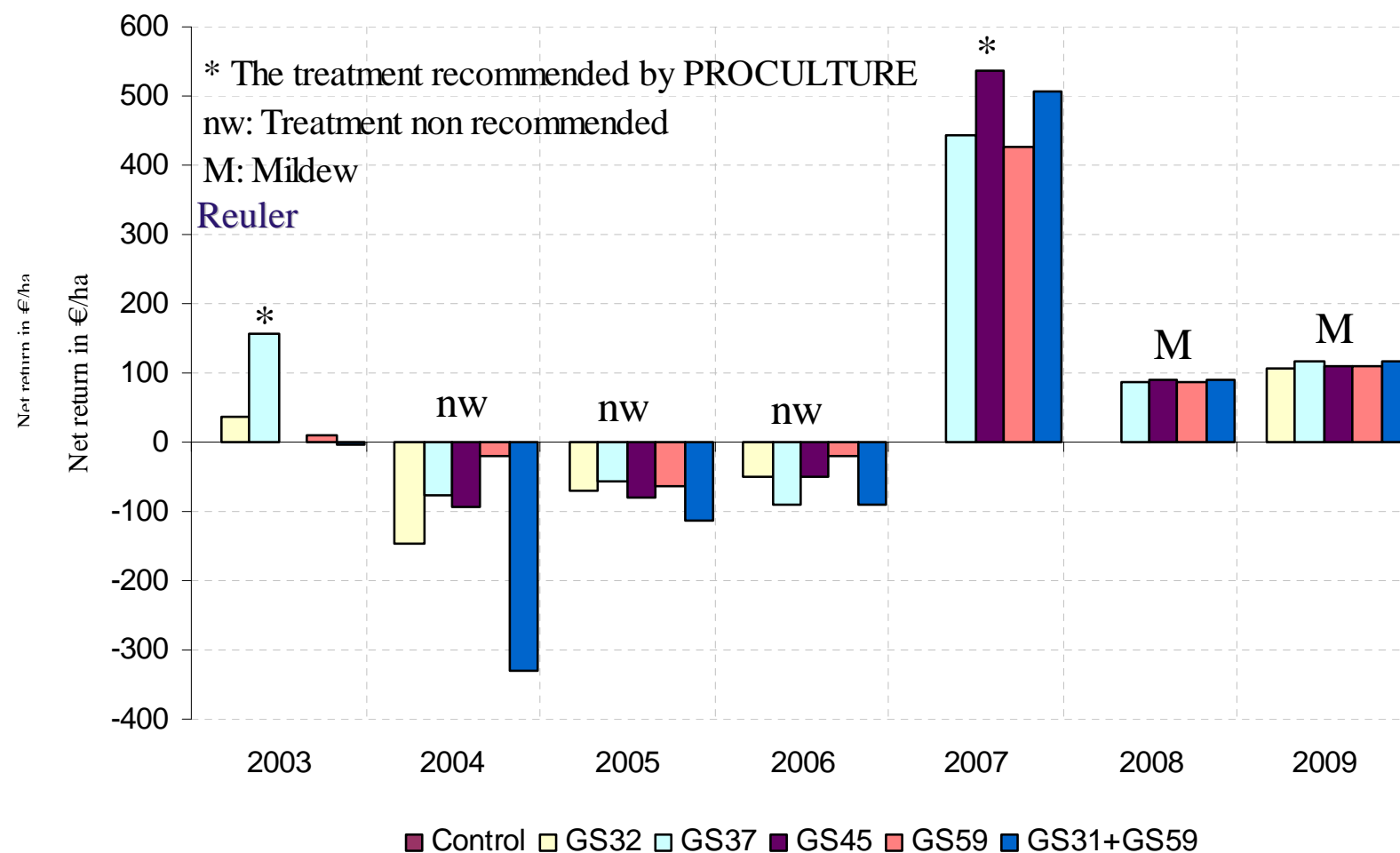
Sowing date →



## Time spray strategies for Septoria Leaf Blotch disease progress on winter wheat: the use of forecasting model



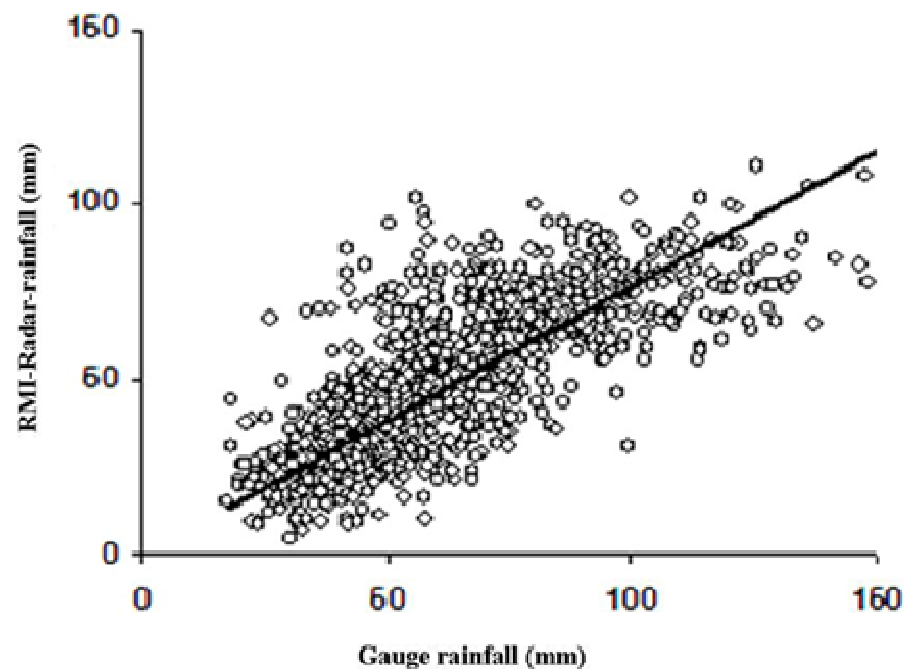
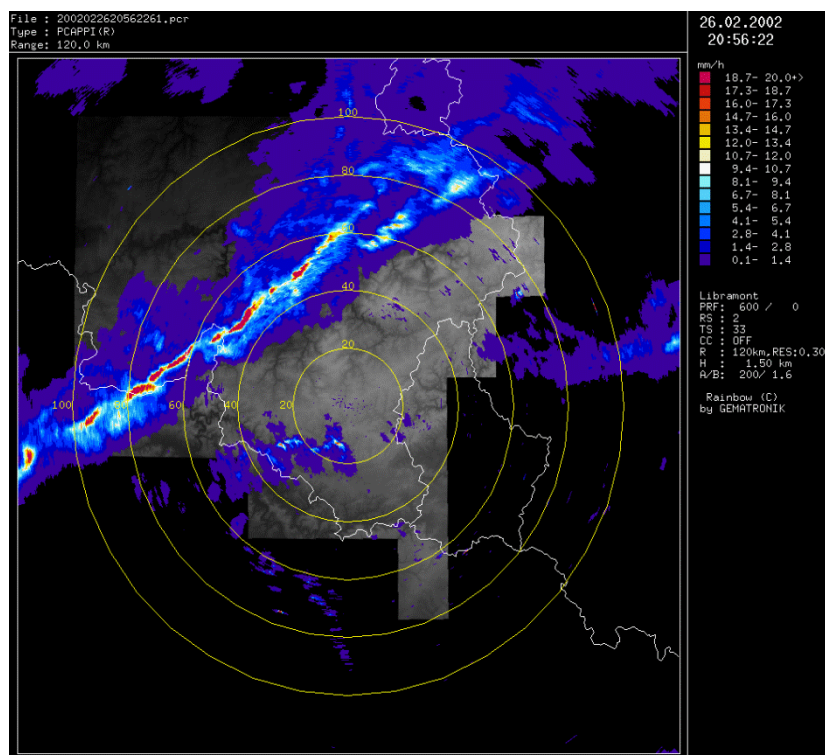
# PROCULTURE Model <sup>3/4</sup>





## Spatial early-warning systems

( $R^2 = 0.75$ ,  $P < 0.001$ )



The use of radar could be a promising alternative for site-specific SLB risk assessment.

# Leaf rust 1/3

Infections occur preferentially at night (De vallavieille-Pope et al. 1995)

Most rapid spread occurs at

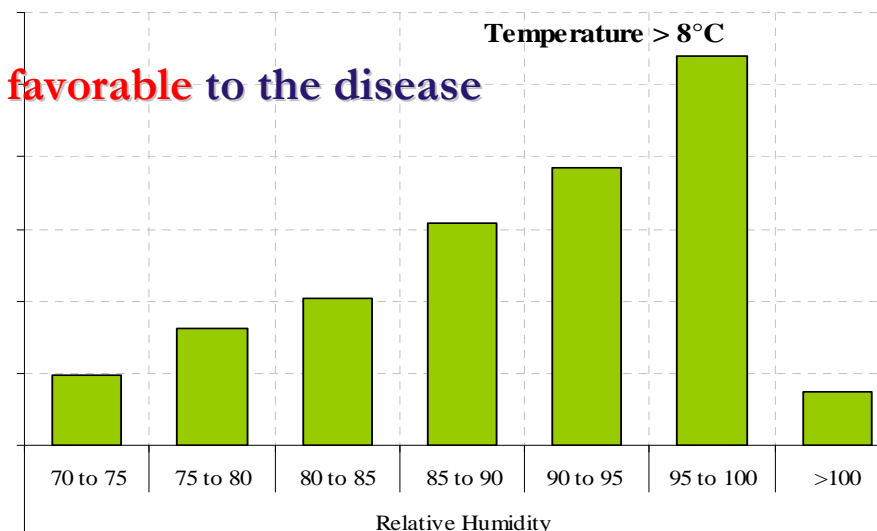
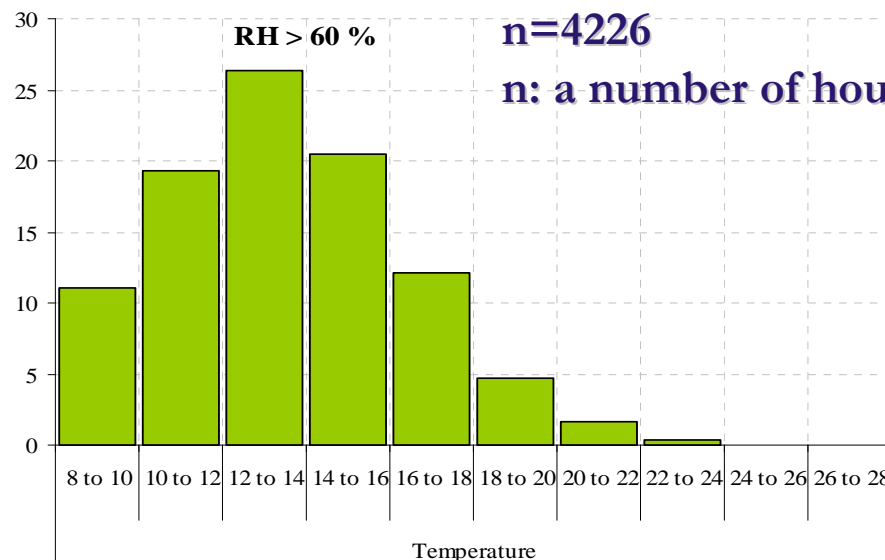
The originality of our work is  
the development of a  
predictive model based on the  
analysis of the night weather  
and leaf rust incidence.

Slow disease development  
in spring – infection by  
wind borne uredospores

and basidiospores.  
Aecidial stage found on  
alternate hosts.

Over-winters on volunteers

# Leaf rust 2/3



- The weather conditions which supported the germination of brown rust between 2000 and 2003 are especially the night temperature classes ranging **between 10 and 18 °C** with an **optimum between 14 and 16°C** associated with ambient relative humidity ranging between **60 and 100%**.
- Night T°C ranging between 24 and 28°C never occurred during the 4 years of the study.
- The frequency of the night temperature classes ranging **between 18 and 24 °C** does not exceed **10%**.

# Leaf rust 3/3

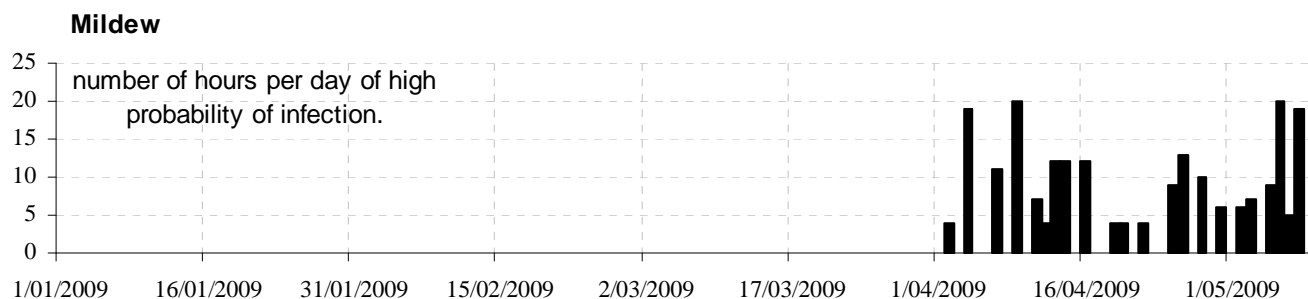
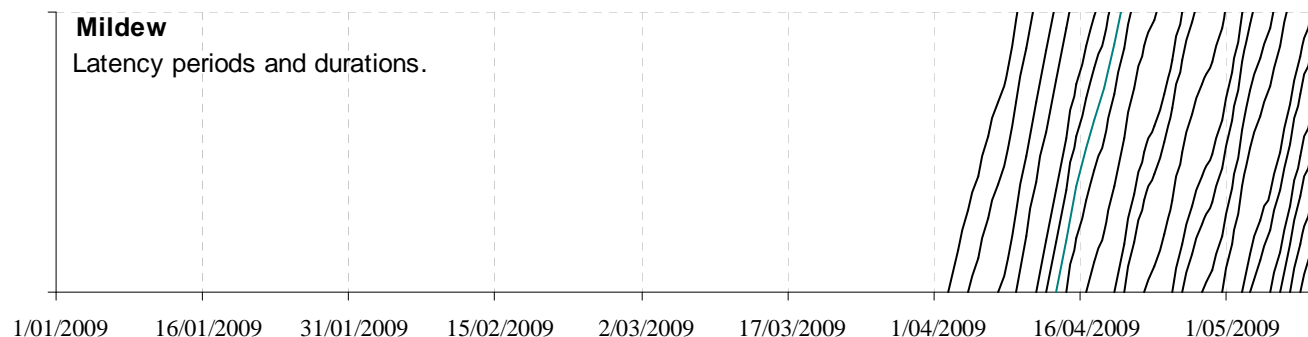




# Wheat Powdery Mildew <sup>1/2</sup>

Two major climatic factors favored the 2003 and 2009 outbreaks, i.e. a daily mean temperature between 15 and 22°C and a relative humidity of at least 80% during April-June

Over 2003 to 2010, a significant difference in severity





# Wheat Powdery Mildew <sup>2/2</sup>

Characterization and modeling of temporal and spatial variability of wetness duration in the context of a national control

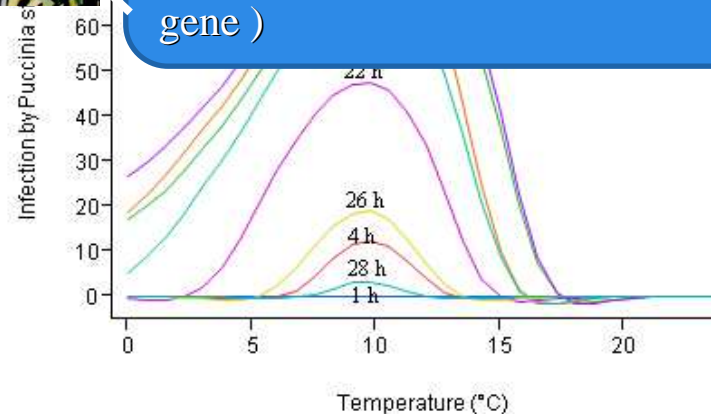


# Yellow rust <sup>1/2</sup>



The minimal continuous wet periods for WSR revealed by the Monte Carlo's method necessary for infection is 4 h at optimal temperature (8 to 15°C).

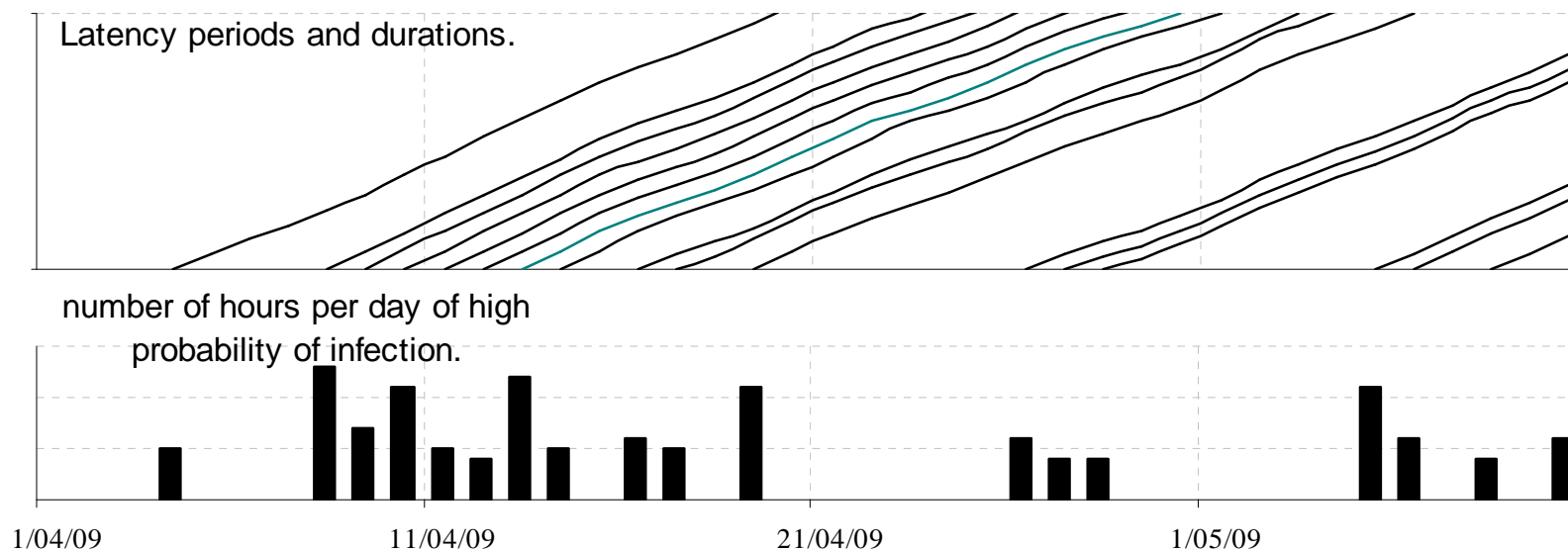
Focus of yellow rust  
Everlange, 2000  
Cultivar: Flair (Yr17 resistance gene)





# Yellow rust <sup>2/2</sup>

Infection parameters: Relative humidity  $\geq 90\%$  for at least 4 hours, temperature between 4 and 16 °C for at least 48 hours and rainfall occurrence (one hour with precipitation  $\geq 0.1$  mm)



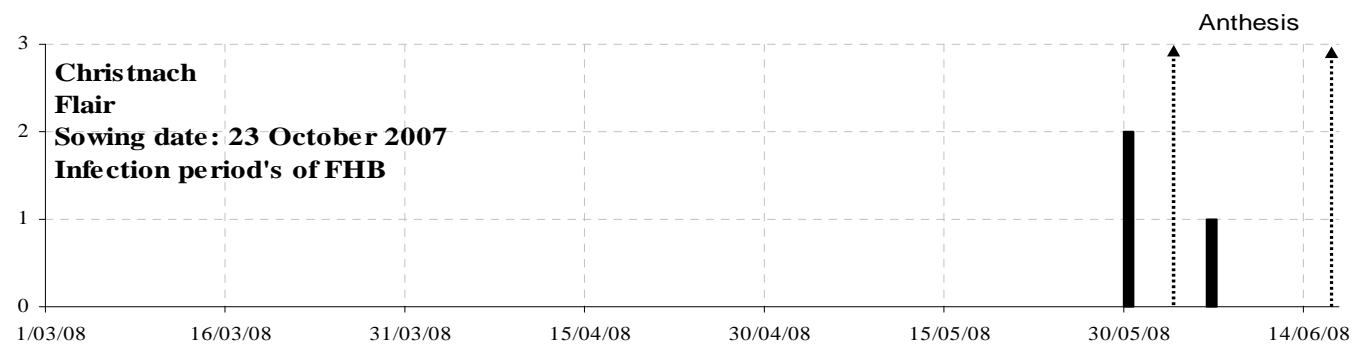
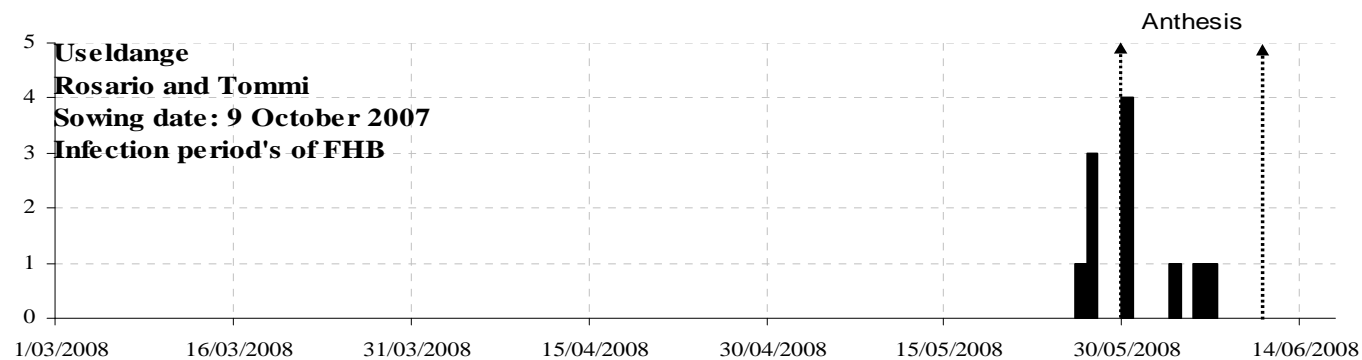
Simulations of infection and latency data by the model are in very good agreement ( $R = 0.92$ ,  $P < 0.05$ ) with the observational data.

# Fusarium head blight 1/2



# Fusarium head blight 2/2

Infection parameters: any event  $\geq 1$  hour during which there is simultaneously a temperature  $\geq 16^\circ \text{C}$ , a relative humidity  $\geq 90\%$  and a rainfall  $\geq 1$  mm during the first hour.





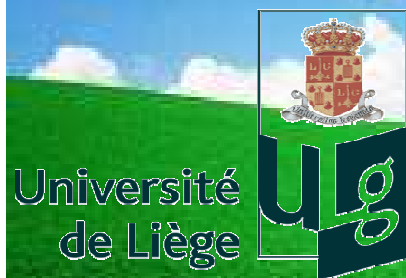
# Conclusions

- ❖ Assessment of the infection periods of SLB by PROCULTURE model in the G-D of Luxembourg achieved an accuracy of 85%.
- ❖ Extension of the disease warning to rusts (WLR and WSR), WPM and FHB.
- ❖ A range of night temperature favourable to the germination of WLR which ranged between 10°C and 18°C with an optimum between 14°C and 16°C.
- ❖ Below this temperature range, the disease progress is nearly stopped. Rainfall is also compulsory but only for initiation of infection, for laying down spores on leaves.
- ❖ WPM appeared much earlier in the northern Öesling (GS 30, pseudostem erection) than in the southern Gutland (GS 39, flag leaf ligule visible)
- ❖ In our study, the minimum continuous wet periods for WSR revealed by Monte Carlo Analysis for an infection is 4 h at optimal temperature (8 to 15°C).

# Perspectives

- ❖ Simplify and possibly map the output of PROCULTURE to make easier the access to information for the farmers  
**Spatial early-warning systems by using weather radar data as the input for the PROCULTURE simulation model**
- ❖ In-depth studies of the microclimatic conditions favourable to the WLR (leaf wetness, RH, T° gradient under the canopy)
  - This future research project will aim at a better understanding of weather parameters that contribute to the development of leaf rust epidemics by studying historical weather and disease incidence data.
  - Another goal is to study inoculum arrival and disease dispersal in the field to better understand and control the WLR in ways that are more respectful of the environment.
  - Study of virulence tests using the European and world differentials and the Yr single gene lines.

# Thank You !



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